

SCORE Search Results Details for Application 10516759 and Search Result 20081112_112530_us-10-516-759-14_copy_24_81.rapbm.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
Page	List	Overview	FAQ	Suggestions

This page gives you Search Results detail for the Application 10516759 and Search Result 20081112_112530_us-10-516-759-14_copy_24_81.rapbm.

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OM protein - protein search, using sw model

Run on: November 12, 2008, 12:17:27 ; Search time 261 Seconds
(without alignments)
214.339 Million cell updates/sec

Title: US-10-516-759-14_COPY_24_81
Perfect score: 350
Sequence: 1 DIKHNRPRRDCVAEGKVCDP.....RNYSRGGVCVTHCNFLNGEP 58

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 4190237 seqs, 964527045 residues

Total number of hits satisfying chosen parameters: 4190237

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA_Main:*

- 1: /ABSS/Data/CRF/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
- 2: /ABSS/Data/CRF/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
- 3: /ABSS/Data/CRF/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
- 4: /ABSS/Data/CRF/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*
- 5: /ABSS/Data/CRF/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
- 6: /ABSS/Data/CRF/ptodata/2/pubpaa/US11A_PUBCOMB.pep:*
- 7: /ABSS/Data/CRF/ptodata/2/pubpaa/US11B_PUBCOMB.pep:*
- 8: /ABSS/Data/CRF/ptodata/2/pubpaa/US12_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	350	100.0	82	5	US-10-516-759-14	Sequence 14, Appl
2	350	100.0	211	6	US-11-443-428A-762461	Sequence 762461,
3	350	100.0	569	6	US-11-043-591-97	Sequence 97, Appl
4	350	100.0	640	5	US-10-516-759-2	Sequence 2, Appli
5	350	100.0	726	6	US-11-443-428A-762452	Sequence 762452,
6	350	100.0	743	6	US-11-443-428A-762450	Sequence 762450,
7	350	100.0	814	6	US-11-443-428A-762451	Sequence 762451,
8	350	100.0	1039	6	US-11-443-428A-759211	Sequence 759211,
9	350	100.0	1276	6	US-11-443-428A-759210	Sequence 759210,
10	350	100.0	1298	6	US-11-365-989-114	Sequence 114, App
11	350	100.0	1298	6	US-11-443-428A-759215	Sequence 759215,
12	350	100.0	1300	6	US-11-043-591-96	Sequence 96, Appl
13	350	100.0	1302	6	US-11-043-591-98	Sequence 98, Appl
14	350	100.0	1342	4	US-10-172-620-16	Sequence 16, Appl
15	350	100.0	1342	4	US-10-207-498-2	Sequence 2, Appli
16	350	100.0	1342	4	US-10-341-434-79	Sequence 79, Appl
17	350	100.0	1342	4	US-10-295-027-1238	Sequence 1238, Ap
18	350	100.0	1342	4	US-10-693-030-4	Sequence 4, Appli
19	350	100.0	1342	5	US-10-723-860-2185	Sequence 2185, Ap
20	350	100.0	1342	5	US-10-482-029-265	Sequence 265, App
21	350	100.0	1342	5	US-10-756-149-5294	Sequence 5294, Ap
22	350	100.0	1342	5	US-10-770-726-63	Sequence 63, Appl
23	350	100.0	1342	5	US-10-219-051B-8640	Sequence 8640, Ap
24	350	100.0	1342	5	US-10-563-888A-2	Sequence 2, Appli
25	350	100.0	1342	5	US-10-503-486-6	Sequence 6, Appli
26	350	100.0	1342	5	US-10-567-867-227	Sequence 227, App
27	350	100.0	1342	5	US-10-533-069-322	Sequence 322, App
28	350	100.0	1342	5	US-10-516-759-1	Sequence 1, Appli
29	350	100.0	1342	6	US-11-037-713-13	Sequence 13, Appl
30	350	100.0	1342	6	US-11-113-202-12	Sequence 12, Appl
31	350	100.0	1342	6	US-11-113-202-14	Sequence 14, Appl
32	350	100.0	1342	6	US-11-406-679-2	Sequence 2, Appli
33	350	100.0	1342	6	US-11-129-740-267	Sequence 267, App
34	350	100.0	1342	6	US-11-443-428A-759208	Sequence 759208,
35	350	100.0	1342	6	US-11-429-090-204	Sequence 204, App
36	350	100.0	1342	6	US-11-582-861-9026	Sequence 9026, Ap
37	350	100.0	1342	6	US-11-591-229-409	Sequence 409, App
38	350	100.0	1342	7	US-11-649-722-390	Sequence 390, App
39	350	100.0	1360	5	US-10-940-774-8022	Sequence 8022, Ap
40	338	96.6	203	6	US-11-443-428A-762456	Sequence 762456,
41	338	96.6	203	6	US-11-443-428A-762460	Sequence 762460,
42	338	96.6	562	4	US-10-159-353B-2	Sequence 2, Appli
43	305	87.1	1339	5	US-10-840-512-214	Sequence 214, App
44	304	86.9	1339	5	US-10-219-051B-8638	Sequence 8638, Ap
45	304	86.9	1339	5	US-10-743-643-631	Sequence 631, App

ALIGNMENTS

RESULT 1

US-10-516-759-14
; Sequence 14, Application US/10516759
; Publication No. US20080057064A1
; GENERAL INFORMATION:
; APPLICANT: ZENSUN(SHANGHAI)SCIENCE AND TECHNOLOGY LIMITED
; APPLICANT: Zhou, Mingdong
; TITLE OF INVENTION: ERBB3 BASED METHODS AND COMPOSITIONS FOR
; TITLE OF INVENTION: TREATING NEOPLASMS
; FILE REFERENCE: 11748-006-999
; CURRENT APPLICATION NUMBER: US/10/516,759
; CURRENT FILING DATE: 2004-12-02
; PRIOR APPLICATION NUMBER: PCT/CN03/00217
; PRIOR FILING DATE: 2003-03-26
; PRIOR APPLICATION NUMBER: CH 02116259
; PRIOR FILING DATE: 2002-03-26
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 82
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-516-759-14

Query Match	100.0%;	Score 350;	DB 5;	Length 82;
Best Local Similarity	100.0%;	Pred. No. 7.4e-27;		
Matches	58;	Conservative	0;	Mismatches 0; Indels 0; Gaps 0;

Qy	1	DIKHNRP	RRDCVA	EKGKVC	DPLCSS	GGCWGP	GPQGCL	SCRNYS	RRGGVC	VTHCNF	LNGEP	58
Db	24	DIKHNRP	RRDCVA	EKGKVC	DPLCSS	GGCWGP	GPQGCL	SCRNYS	RRGGVC	VTHCNF	LNGEP	81

RESULT 2

US-11-443-428A-762461
; Sequence 762461, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31

Query Match 100.0%; Score 350; DB 6; Length 211;
Best Local Similarity 100.0%; Pred. No. 1.7e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 DIKHNRRPRDCVAEGKVCDPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
 |||||||

Db 124 DIKHNRRPRDCVAEGKVCDPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 181

US-11-043-591-97

; Publication No. US20070082337A1

; GENERAL INFORMATION:

```
; APPLICANT: Sorek, Rotem
; APPLICANT: Pollock, Sarah
; APPLICANT: Diber, Alex
; APPLICANT: Levine, Zurit
; APPLICANT: Nemzer, Sergey
; APPLICANT: Kol, Guy
; APPLICANT: Wool, Assaf
; APPLICANT: Haviv, Ami
; APPLICANT: Cohen, Yuval
; APPLICANT: Cohen, Yossi
; APPLICANT: Shemesh, Ronen
; APPLICANT: Savitsky, Kinneret
```

; TITLE OF INVENTION: METHODS OF IDENTIFYING PUTATIVE GENE PRODUCTS BY INTERSPECIES
SEQUENCE

; TITLE OF INVENTION: COMPARISON AND BIOMOLECULAR SEQUENCES UNCOVERED THEREBY

; FILE REFERENCE: 28486

; CURRENT APPLICATION NUMBER: US/11/043,591

; CURRENT FILING DATE: 2005-01-27

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; NUMBER OF SEQ ID NOS: 469
```

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; SOFTWARE: PatentIn version 3.2
```

; SEQ ID NO 97

; LENGTH: 569

```
; TYPE: PRT
```

```
; ORGANISM: Artificial sequence
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; FEATURE:
```

; OTHER INFORMATION: A novel predicted alternative spliced variant protein product

US-11-043-591-97

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Query Match      100.0%;  Score 350;  DB 6;  Length 569;
Best Local Similarity 100.0%;  Pred. No. 3.9e-26;
Matches    58;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;
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```
Qy          1 DIKHNRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
            |||
Db          483 DIKHNRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 540
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RESULT 4

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US-10-516-759-2
; Sequence 2, Application US/10516759
; Publication No. US20080057064A1
; GENERAL INFORMATION:
; APPLICANT: ZENSUN(SHANGHAI)SCIENCE AND TECHNOLOGY LIMITED
; APPLICANT: Zhou, Mingdong
; TITLE OF INVENTION: ERBB3 BASED METHODS AND COMPOSITIONS FOR
; TITLE OF INVENTION: TREATING NEOPLASMS
; FILE REFERENCE: 11748-006-999
; CURRENT APPLICATION NUMBER: US/10/516,759
; CURRENT FILING DATE: 2004-12-02
; PRIOR APPLICATION NUMBER: PCT/CN03/00217
; PRIOR FILING DATE: 2003-03-26
; PRIOR APPLICATION NUMBER: CH 02116259
; PRIOR FILING DATE: 2002-03-26
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 640
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-516-759-2
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Query Match          100.0%; Score 350; DB 5; Length 640;
Best Local Similarity 100.0%; Pred. No. 4.3e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy          1 DIKHNRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
            |||
Db          483 DIKHNRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 540
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RESULT 5

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US-11-443-428A-762452
; Sequence 762452, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
```

```
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 762452
; LENGTH: 726
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-443-428A-762452
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Query Match 100.0%; Score 350; DB 6; Length 726;
Best Local Similarity 100.0%; Pred. No. 4.8e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 DIKHNRRPRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
 |||
 Db 124 DIKHNRRPRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 181

RESULT 6

US-11-443-428A-762450

; Sequence 762450, Application US/11443428A

; Publication No. US20070083334A1

; GENERAL INFORMATION:

```
; APPLICANT: Mintz, Liat
```

; APPLICANT: Xie, Hanqing

```
; APPLICANT:  Dahari, Dvir
```

; APPLICANT: Levanon, Erez

; APPLICANT: Freilich, Shiri

; APPLICANT: Beck, Nili

; APPLICANT: Zhu, Wei-Yong

; APPLICANT: Wasserman, Alon

; APPLICANT: Hermesh, Chen

; APPLICANT: Azar, Idit

; APPLICANT: Bernstein, Jeanne

; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES

; FILE REFERENCE: 02/23929

; CURRENT APPLICATION NUMBER: US/11/443,428A

; CURRENT FILING DATE: 2006-05-31

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; NUMBER OF SEQ ID NOS: 1034312
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; SOFTWARE: PatentIn version 3.1
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; SEQ ID NO 762450

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; LENGTH: 743
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; TYPE: PRT
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; ORGANISM: Homo sapiens

US-11-443-428A-762450

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Query Match      100.0%;  Score 350;  DB 6;  Length 743;
Best Local Similarity 100.0%;  Pred. No. 4.9e-26;
Matches    58;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;
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Qy 1 DIKHNRRPRDCVAEGKVCDPLCSSGGCWGP GPGQCLSCRNYSRGGVCVTHCNFLNGEP 58

Db 124 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 181

RESULT 7
US-11-443-428A-762451
; Sequence 762451, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 762451
; LENGTH: 814
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-443-428A-762451

Query Match 100.0%; Score 350; DB 6; Length 814;
Best Local Similarity 100.0%; Pred. No. 5.3e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
Db 124 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 181

RESULT 8
US-11-443-428A-759211
; Sequence 759211, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon

```

Query Match          100.0%;   Score 350;   DB 6;   Length 1039;
Best Local Similarity 100.0%;   Pred. No. 6.5e-26;
Matches    58;   Conservative    0;   Mismatches    0;   Indels    0;   Gaps    0;

Qy          1 DIKHNRRPRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
            |||
Db          180 DIKHNRRPRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 237

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US-11-443-428A-759210

Query Match	100.0%;	Score 350;	DB 6;	Length 1276;
Best Local Similarity	100.0%;	Pred. No. 7.8e-26;		


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Matches      58;  Conservative      0;  Mismatches      0;  Indels      0;  Gaps      0;

Qy           1 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
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Db          417 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 474

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RESULT 10

US-11-365-989-114

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; Sequence 114, Application US/11365989
; Publication No. US20060199226A1
; GENERAL INFORMATION:
; APPLICANT: Schiffer, Hans
; TITLE OF INVENTION: FUNCTIONAL BIOLUMINESCENCE ENERGY
; TITLE OF INVENTION: RESONANCE TRANSFER (BRET) ASSAY TO SCREEN, IDENTIFY AND
; TITLE OF INVENTION: CHARACTERIZE RECEPTOR TYROSINE KINASE LIGANDS
; FILE REFERENCE: ACADIA.072A
; CURRENT APPLICATION NUMBER: US/11/365,989
; CURRENT FILING DATE: 2006-03-01
; PRIOR APPLICATION NUMBER: 60/658,319
; PRIOR FILING DATE: 2005-03-02
; NUMBER OF SEQ ID NOS: 234
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 114
; LENGTH: 1298
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Amino acid sequence of HER3

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US-11-365-989-114

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Query Match      100.0%;  Score 350;  DB 6;  Length 1298;
Best Local Similarity 100.0%;  Pred. No. 7.9e-26;
Matches      58;  Conservative      0;  Mismatches      0;  Indels      0;  Gaps      0;

Qy           1 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
             |||
Db          439 DIKHNRPRRDCVAEGKVC DPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 496

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RESULT 11

US-11-443-428A-759215

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; Sequence 759215, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen

```

```
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 759215
;   LENGTH: 1298
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-443-428A-759215
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Query Match 100.0%; Score 350; DB 6; Length 1298;
Best Local Similarity 100.0%; Pred. No. 7.9e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 DIKHNRRPRDCVAEGKVCPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 58
 |||
 Db 439 DIKHNRRPRDCVAEGKVCPLCSSGGCWGPGPGQCLSCRNYSRGGVCVTHCNFLNGEP 496

RESULT 12

US-11-043-591-96

```

; Sequence 96, Application US/11043591
; Publication No. US20070082337A1
; GENERAL INFORMATION:
; APPLICANT: Sorek, Rotem
; APPLICANT: Pollock, Sarah
; APPLICANT: Diber, Alex
; APPLICANT: Levine, Zurit
; APPLICANT: Nemzer, Sergey
; APPLICANT: Kol, Guy
; APPLICANT: Wool, Assaf
; APPLICANT: Haviv, Ami
; APPLICANT: Cohen, Yuval
; APPLICANT: Cohen, Yossi
; APPLICANT: Shemesh, Ronen
; APPLICANT: Savitsky, Kinneret
; TITLE OF INVENTION: METHODS OF IDENTIFYING PUTATIVE GENE PRODUCTS BY INTERSPECIES
SEQUENCE
; TITLE OF INVENTION: COMPARISON AND BIOMOLECULAR SEQUENCES UNCOVERED THEREBY
; FILE REFERENCE: 28486
; CURRENT APPLICATION NUMBER: US/11/043,591
; CURRENT FILING DATE: 2005-01-27
; NUMBER OF SEQ ID NOS: 469
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 96
; LENGTH: 1300
; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: A novel predicted alternative spliced variant protein product

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Best Local Similarity 100.0%; Pred. No. 8.1e-26;
Matches 58; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	DIKHNRPRRDCVAEGKVC	DPLCSSGGCWGP	PGQCLSCRNYSRGGVCVTHCNFLNGEP	58
Db	483	DIKHNRPRRDCVAEGKVC	DPLCSSGGCWGP	PGQCLSCRNYSRGGVCVTHCNFLNGEP	540

Search completed: November 12, 2008, 12:21:51
Job time : 264 secs

SCORE 3.0